

1/4

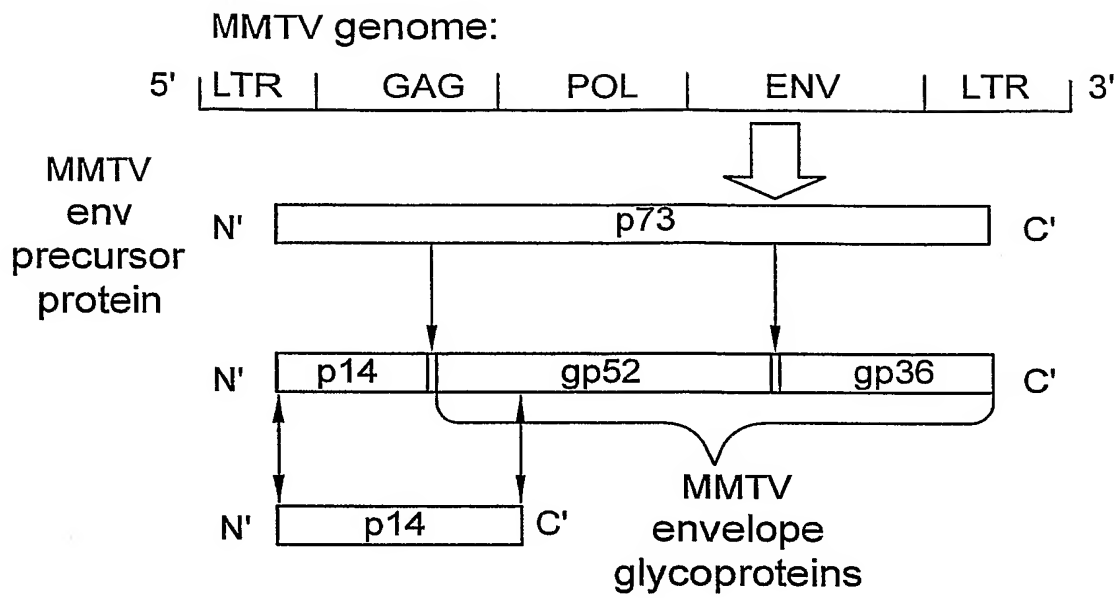


FIG. 1A

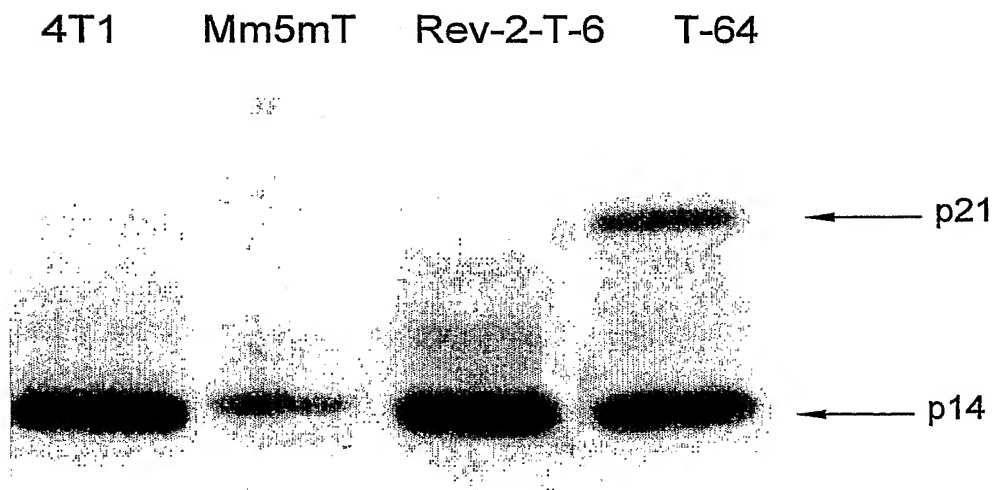
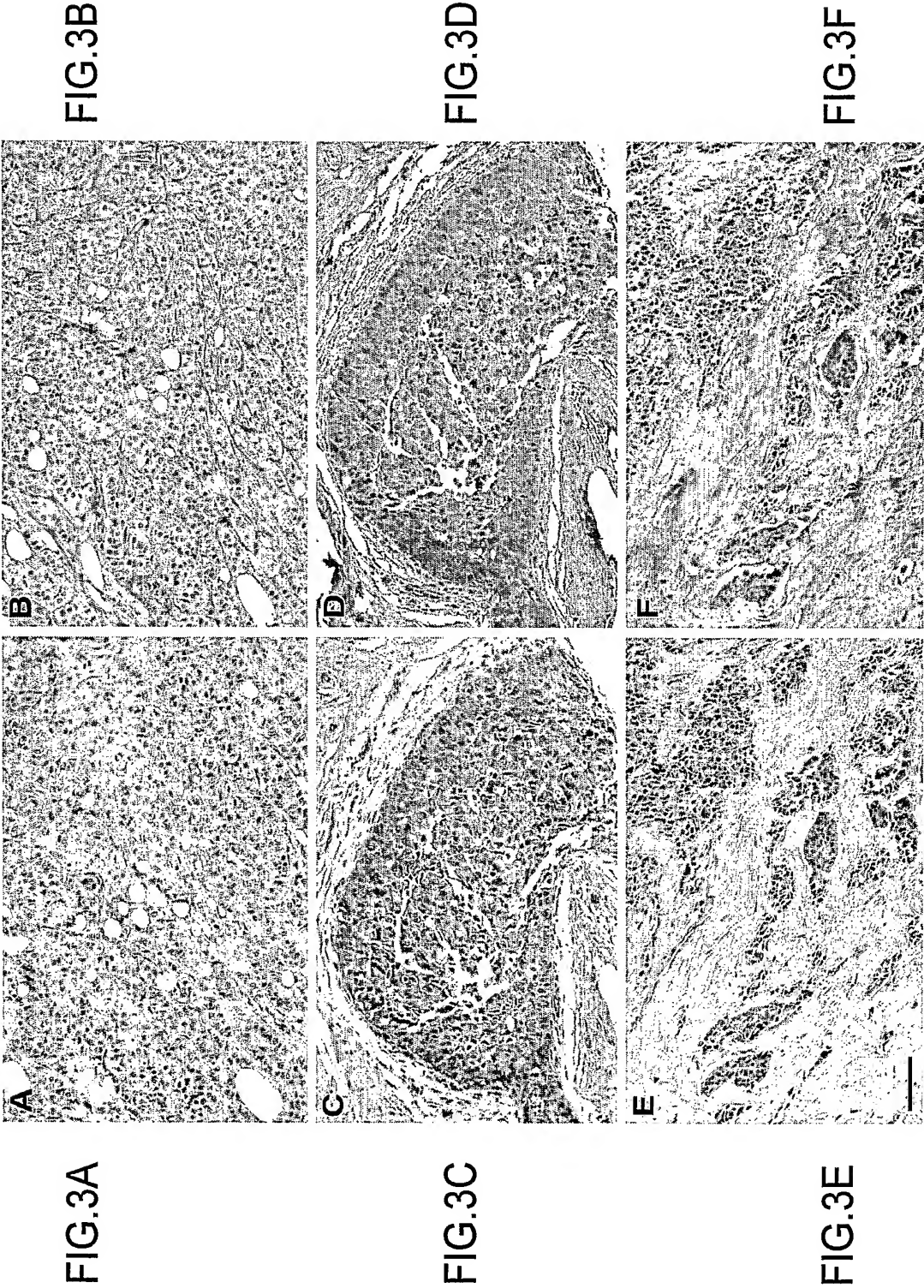


FIG. 1B

af248270 2 GSLTDSSDLSL SGKDRRPRLALRRKRRREM RKITKKVRRMTLDPIKEKTA 51
||| ||| ||| | . | | | | | | | | : | | | | | | | |
p14 7 GSPTGSSDLLL SGKKQRPHLALRRKRRREM KKNRKVRRMNLDLIKEKTA 56
52 WQQLQALISEAEEVLKTSQTPQTSLTLFLALLSVLGPLPVTG 93
|| | | | | | | | | | | | | | | | | | | | | | | | |
57 WQHLQALISEAEEVLKTSQTPQTSLTLFLALLSVLGPPPPVTG 98

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FIG. 2



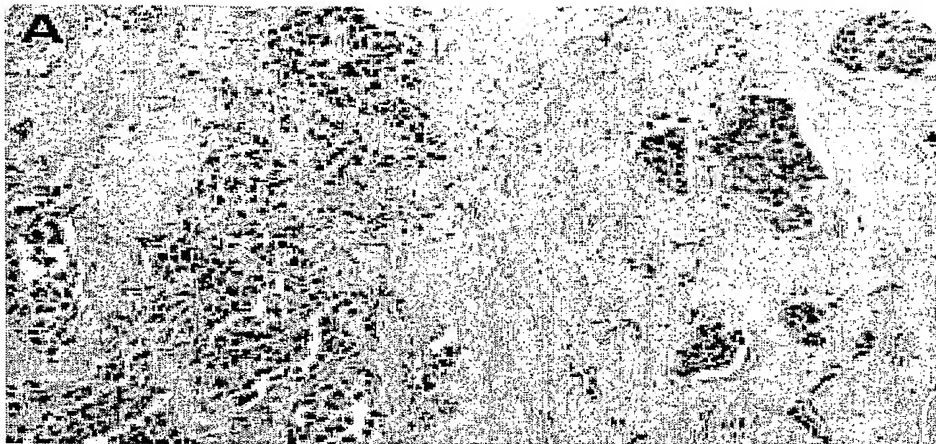


FIG. 4A

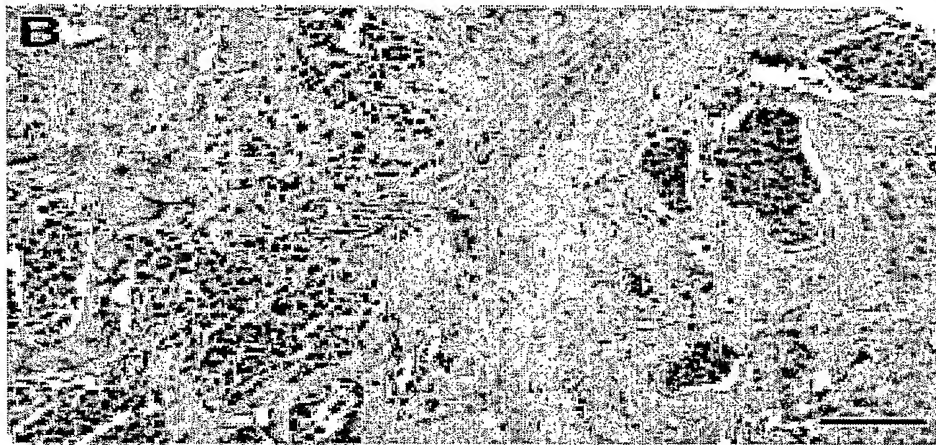


FIG. 4B



FIG. 4C